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SEQUENCE LISTING

<110> PIERRE FABRE MEDICAMENT

<120> USE OF BACTERIAL MEMBRANE FRACTIONS WITH
IMMUNOSTIMULANT ACTIVITY IN THE TREATMENT OF
CANCERS, METHODS FOR PREPARING THEM AND THE
PHARMACEUTICAL COMPOSITIONS CONTAINING THEM

<130> D17974

<140>

<141>

<160> 2

<170> PatentIn Vers. 2.0

<210> 1

<211> 1035

<212> DNA

<213> Klebsiella pneumoniae

<220>

<221> exon

<222> (1)..(1032)

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<221> intron

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tat gca ggt ggt aaa ctg ggt tgg tcc cag tat cac gac acc ggt ttc 96

Tyr Ala Gly Gly Lys Leu Gly Trp Ser Gln Tyr His Asp Thr Gly Phe

20 25 30

tac ggt aac ggt ttc cag aac aac ggt ccg acc cgt aac gat cag 144

Tyr Gly Asn Gly Phe Gln Asn Asn Asn Gly Pro Thr Arg Asn Asp Gln

35 40 45

ctt ggt gct ggt gcg ttc ggt ggt tac cag gtt aac ccg tac ctc ggt 192

Leu Gly Ala Gly Ala Phe Gly Gly Tyr Gln Val Asn Pro Tyr Leu Gly

50 55 60

ttc gaa atg ggt tat gac tgg ctg ggc cgt atg gca tat aaa ggc agc 240

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Phe	Glu	Met	Gly	Tyr	Asp	Trp	Leu	Gly	Arg	Met	Ala	Tyr	Lys	Gly	Ser		
65															80		
gtt	gac	aac	ggt	gct	tcc	aaa	gct	cag	ggc	gtt	cag	ctg	acc	gct	aaa	288	
Val	Asp	Asn	Gly	Ala	Phe	Lys	Ala	Gln	Gly	Val	Gln	Leu	Thr	Ala	Lys		
85															95		
ctg	ggt	tac	ccg	atc	act	gac	gat	ctg	gac	atc	tac	acc	cgt	ctg	ggc	336	
Leu	Gly	Tyr	Pro	Ile	Thr	Asp	Asp	Leu	Asp	Ile	Tyr	Thr	Arg	Leu	Gly		
100															110		
ggc	atg	gtt	tgg	ccg	gct	gac	tcc	aaa	ggc	aac	tac	gct	tct	acc	ggc	384	
Gly	Met	Val	Trp	Arg	Ala	Asp	Ser	Lys	Gly	Asn	Tyr	Ala	Ser	Thr	Gly		
115															125		
gtt	tcc	cgt	agc	gaa	cac	gac	act	ggc	gtt	tcc	cca	gta	ttt	gct	ggc	432	
Val	Ser	Arg	Ser	Glu	His	Asp	Thr	Gly	Val	Ser	Pro	Val	Phe	Ala	Gly		
130															140		
ggc	gta	gag	tgg	gct	gtt	act	cgt	gac	atc	gct	acc	cgt	ctg	gaa	tac	480	
Gly	Val	Glu	Trp	Ala	Val	Thr	Arg	Asp	Ile	Ala	Thr	Arg	Leu	Glu	Tyr		
145															155	160	
cag	tgg	gtt	aac	aac	atc	ggc	gac	ggc	act	gtg	ggt	acc	cgt	cct		528	
Gln	Trp	Val	Asn	Asn	Ile	Gly	Asp	Ala	Gly	Thr	Val	Gly	Thr	Arg	Pro		
165															170	175	
gat	aac	ggc	atg	ctg	agc	ctg	ggc	gtt	tcc	tac	cgc	tcc	ggt	cag	gaa	576	
Asp	Asn	Gly	Met	Leu	Ser	Leu	Gly	Val	Ser	Tyr	Arg	Phe	Gly	Gln	Glu		
180															185	190	
gat	gct	gca	ccg	gtt	gtt	gct	ccg	gct	ccg	gct	ccg	gaa	gtg			624	
Asp	Ala	Ala	Pro	Val	Val	Ala	Pro	Ala	Pro	Ala	Pro	Ala	Pro	Glu	Val		
195															200	205	
gtt	acc	aag	cac	tcc	acc	ctg	aag	tct	gac	gtt	ctg	tcc	aac	tcc	aac	672	
Ala	Thr	Lys	His	Phe	Thr	Leu	Lys	Ser	Asp	Val	Leu	Phe	Asn	Phe	Asn		
210															215	220	
aaa	gct	acc	ctg	aaa	ccg	gaa	ggt	cag	cag	gct	ctg	gat	cag	ctg	tac	720	
Lys	Ala	Thr	Leu	Lys	Pro	Glu	Gly	Gln	Gln	Ala	Leu	Asp	Gln	Leu	Tyr		
225															230	235	240
act	cag	ctg	agc	aac	atg	gat	ccg	aaa	gac	ggt	tcc	gct	gtt	gtt	ctg	768	
Thr	Gln	Leu	Ser	Asn	Met	Asp	Pro	Lys	Asp	Gly	Ser	Ala	Val	Val	Leu		
245															250	255	
ggc	tac	acc	gac	ccg	atc	ggt	tcc	gaa	gct	tac	aac	cag	cag	ctg	tct	816	
Gly	Tyr	Thr	Asp	Arg	Ile	Gly	Ser	Glu	Ala	Tyr	Asn	Gln	Gln	Leu	Ser		
260															265	270	
gag	aaa	cgt	gct	cag	tcc	gtt	gac	tac	ctg	gtt	gct	aaa	ggc	atc		864	

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Glu Lys Arg Ala Gln Ser Val Val Asp Tyr Leu Val Ala Lys Gly Ile			
275	280	285	
cgg gct ggc aaa atc tcc gct cgc ggc atg ggt gaa tcc aac ccg gtt			912
Pro Ala Gly Lys Ile Ser Ala Arg Gly Met Gly Ser Asn Pro Val			
290	295	300	
act ggc aac acc tgt gac aac gtg aaa gct cgc gct gcc ctg atc gat			960
Thr Gly Asn Thr Cys Asp Asn Val Lys Ala Arg Ala Ala Leu Ile Asp			
305	310	315	320
tgc ctg gct ccg gat cgt cgt gta gag atc gaa gtt aaa ggc tac aaa			1008
Cys Leu Ala Pro Asp Arg Arg Val Glu Ile Glu Val Lys Gly Tyr Lys			
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gaa gtt gta act cag ccg gcg ggt taa			1035
Glu Val Val Thr Gln Pro Ala Gly			
340			

<210> 2

<211> 344

<212> PRT

<213> Klebsiella pneumoniae

<400> 2

Met Lys Ala Ile Phe Val Leu Asn Ala Ala Pro Lys Asp Asn Thr Trp			
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Tyr Ala Gly Gly Lys Leu Gly Trp Ser Gln Tyr His Asp Thr Gly Phe			
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Tyr Gly Asn Gly Phe Gln Asn Asn Asn Gly Pro Thr Arg Asn Asp Gln			
35	40	45	
Leu Gly Ala Gly Ala Phe Gly Gly Tyr Gln Val Asn Pro Tyr Leu Gly			
50	55	60	
Phe Glu Met Gly Tyr Asp Trp Leu Gly Arg Met Ala Tyr Lys Gly Ser			
65	70	75	80
Val Asp Asn Gly Ala Phe Lys Ala Gln Gly Val Gln Leu Thr Ala Lys			
85	90	95	
Leu Gly Tyr Pro Ile Thr Asp Asp Leu Asp Ile Tyr Thr Arg Leu Gly			
100	105	110	
Gly Met Val Trp Arg Ala Asp Ser Lys Gly Asn Tyr Ala Ser Thr Gly			
115	120	125	
Val Ser Arg Ser Glu His Asp Thr Gly Val Ser Pro Val Phe Ala Gly			
130	135	140	

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Gly Val Glu Trp Ala Val Thr Arg Asp Ile Ala Thr Arg Leu Glu Tyr			
145	150	155	160
Gln Trp Val Asn Asn Ile Gly Asp Ala Gly Thr Val Gly Thr Arg Pro			
165	170	175	
Asp Asn Gly Met Leu Ser Leu Gly Val Ser Tyr Arg Phe Gly Gln Glu			
180	185	190	
Asp Ala Ala Pro Val Val Ala Pro Ala Pro Ala Pro Glu Val			
195	200	205	
Ala Thr Lys His Phe Thr Leu Lys Ser Asp Val Leu Phe Asn Phe Asn			
210	215	220	
Lys Ala Thr Leu Lys Pro Glu Gly Gln Gln Ala Leu Asp Gln Leu Tyr			
225	230	235	240
Thr Gln Leu Ser Asn Met Asp Pro Lys Asp Gly Ser Ala Val Val Leu			
245	250	255	
Gly Tyr Thr Asp Arg Ile Gly Ser Glu Ala Tyr Asn Gln Gln Leu Ser			
260	265	270	
Glu Lys Arg Ala Gln Ser Val Val Asp Tyr Leu Val Ala Lys Gly Ile			
275	280	285	
Pro Ala Gly Lys Ile Ser Ala Arg Gly Met Gly Glu Ser Asn Pro Val			
290	295	300	
Thr Gly Asn Thr Cys Asp Asn Val Lys Ala Arg Ala Ala Leu Ile Asp			
305	310	315	320
Cys Leu Ala Pro Asp Arg Arg Val Glu Ile Glu Val Lys Gly Tyr Lys			
325	330	335	
Glu Val Val Thr Gln Pro Ala Gly			
340			